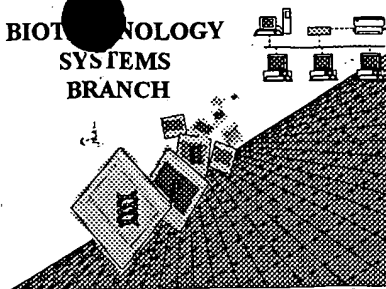


RAW SEQUENCE LISTING **ERROR REPORT**

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/914,176

Source: PCT/09

Date Processed by STIC: 9/13/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

PCT09

RAW SEQUENCE LISTING

DATE: 09/13/2001

PATENT APPLICATION: US/09/914,176

TIME: 17:24:15

Input Set : A:\Wpa.app

Output Set: N:\CRF3\09132001\I914176.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: French, Gary L
 4 Anthony, Richard M
 5 Brown, Timothy J
 7 <120> TITLE OF INVENTION: Identification of Bacteria by Amplification and Probing
 9 <130> FILE REFERENCE: WPA
 11 <140> CURRENT APPLICATION NUMBER: US/09/914,176
 12 <141> CURRENT FILING DATE: 2001-08-24
 14 <150> PRIOR APPLICATION NUMBER: PCT/GB00/00740
 15 <151> PRIOR FILING DATE: 2000-03-01
 17 <150> PRIOR APPLICATION NUMBER: GB 9904804.3
 18 <151> PRIOR FILING DATE: 1999-03-02
 20 <160> NUMBER OF SEQ ID NOS: 63
 22 <170> SOFTWARE: PatentIn Ver. 2.0

ERRORED SEQUENCES

567 <210> SEQ ID NO: 49
 568 <211> LENGTH: 341 21
 569 <212> TYPE: DNA
 570 <213> ORGANISM: Artificial Sequence
 572 <220> FEATURE:
 573 <223> OTHER INFORMATION: Description of Artificial Sequence: primer
 575 <400> SEQUENCE: 49
 E--> 576 gtctggaaag tccgacggt a c 21

FWS →

Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/914,176

DATE: 09/13/2001

TIME: 17:24:16

Input Set : A:\Wpa.app

Output Set: N:\CRF3\09132001\I914176.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number

L:48 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:576 M:252 E: No. of Seq. differs, <211>LENGTH:Input:341 Found:21 SEQ:49